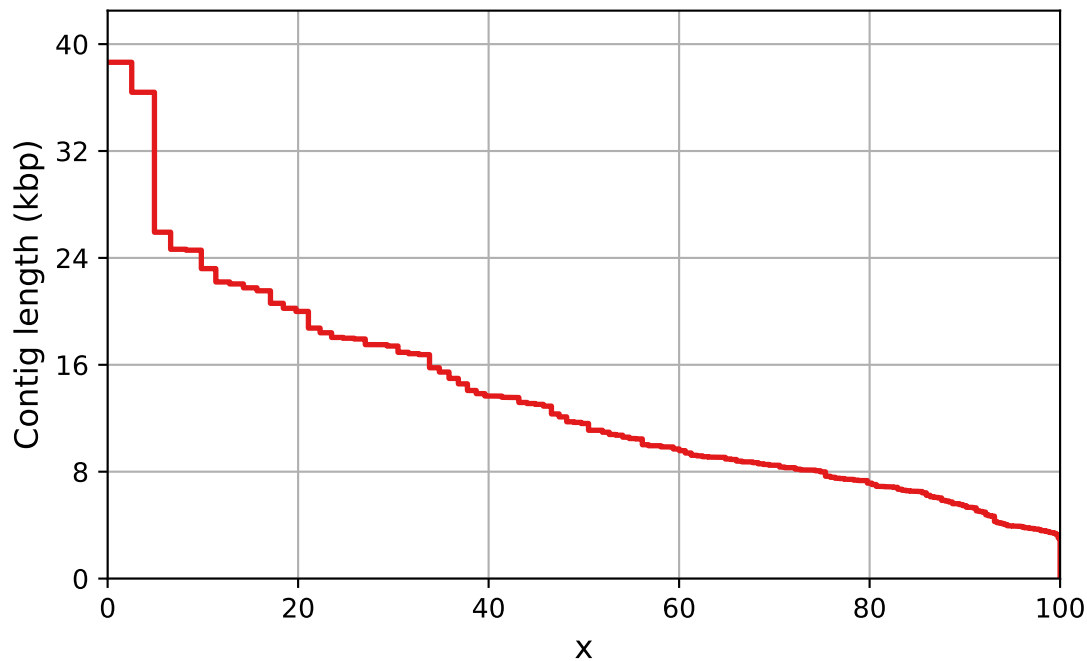


Report

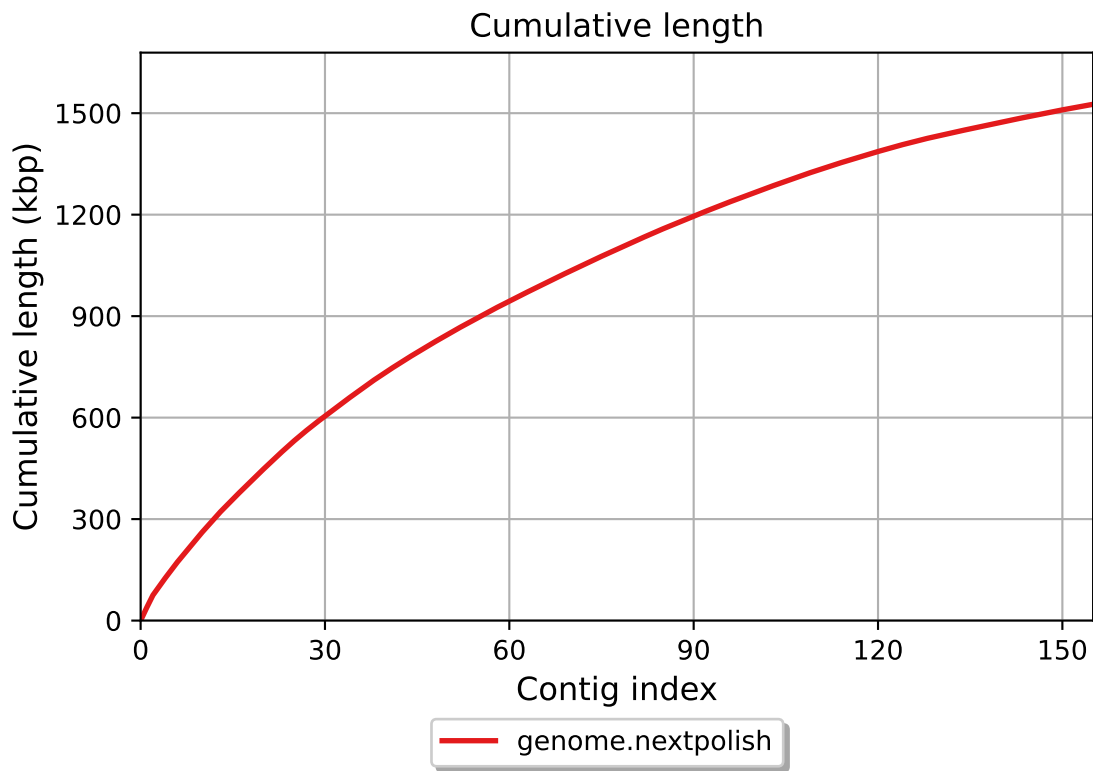
	genome.nextpolish
# contigs (>= 0 bp)	155
# contigs (>= 1000 bp)	155
# contigs (>= 5000 bp)	123
# contigs (>= 10000 bp)	52
# contigs (>= 25000 bp)	3
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1526315
Total length (>= 1000 bp)	1526315
Total length (>= 5000 bp)	1402037
Total length (>= 10000 bp)	866862
Total length (>= 25000 bp)	100959
Total length (>= 50000 bp)	0
# contigs	155
Largest contig	38641
Total length	1526315
GC (%)	57.88
N50	11607
N90	5459
auN	13726.5
L50	43
L90	118
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 1 part

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

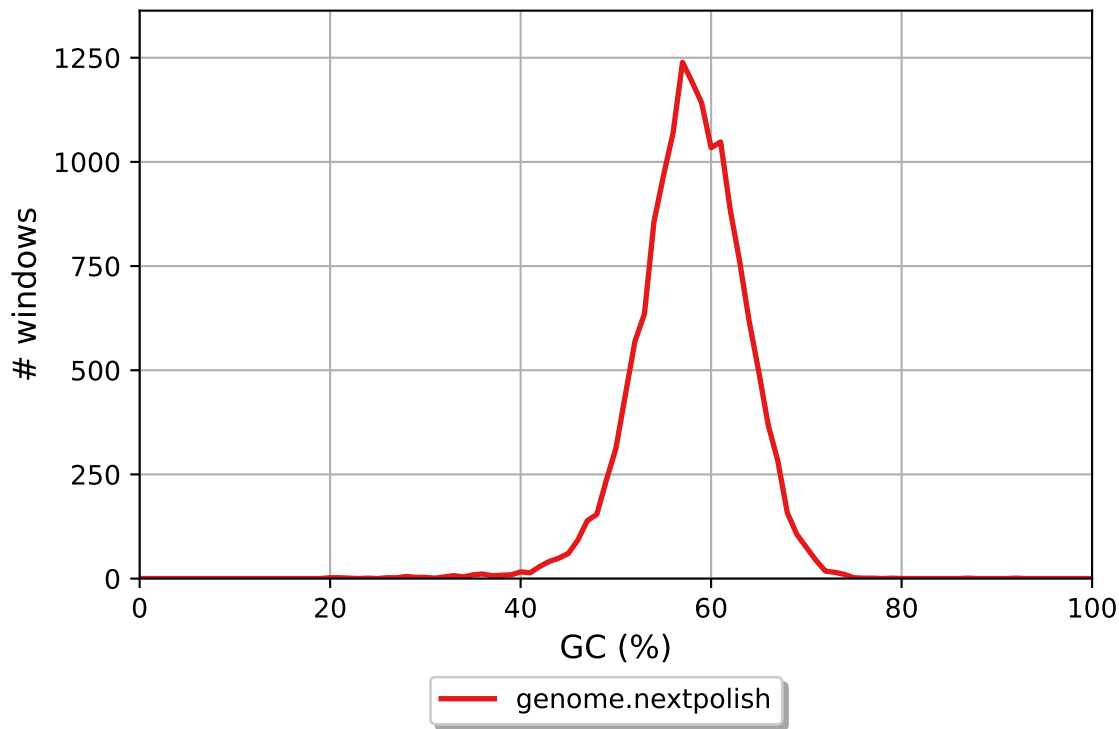
Nx



genome.nextpolish



GC content



genome.nextpolish GC content

